The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/725, 284
Source:	T+WO.
Date Processed by STIC:	10-18-04

# ENTERED



**IFWO** 

RAW SEQUENCE LISTING

DATE: 10/18/2004

PATENT APPLICATION: US/10/725,284

TIME: 09:54:28

Input Set : A:\78003681.app

```
3 <110> APPLICANT: ADLER, JON ELLIOT
             LI, XIAODONG
     5
              STASZEWSKI, LENA
             O'CONNELL, SHAWN
     .6
              ZOZULYA, SERGEY
     7
     9 <120> TITLE OF INVENTION: T1R TASTE RECEPTORS AND GENES ENCODING SAME
     11 <130> FILE REFERENCE: 078003-0280681
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/725,284
C--> 14 <141> CURRENT FILING DATE: 2003-12-02
     16 <150> PRIOR APPLICATION NUMBER: 60/259,227
     17 <151> PRIOR FILING DATE: 2001-01-03
     19 <150> PRIOR APPLICATION NUMBER: 60/284,547
     20 <151> PRIOR FILING DATE: 2001-04-19
     22 <160> NUMBER OF SEQ ID NOS: 24
     24 <170> SOFTWARE: PatentIn Ver. 2.1
     26 <210> SEQ ID NO: 1
     27 <211> LENGTH: 876
     28 <212> TYPE: DNA
     29 <213> ORGANISM: Homo sapiens
     31 <400> SEQUENCE: 1
     32 agectggcag tggcctcagg cagagtctga cgcgcacaaa ctttcaggcc caggaagcga 60
     33 ggacaccact ggggccccag ggtgtggcaa gtgaggatgg caagggtttt gctaaacaaa 120
     34 teetetgeee geteeeegee eegggeteae teeatgtgag geeceagteg gggeageeae 180
     35 ctgccgtgcc tgttggaagt tgcctctgcc atgctgggcc ctgctgtcct gggcctcagc 240
     36 ctctgggctc tcctgcaccc tgggacgggg gccccattgt gcctgtcaca gcaacttagg 300
     37 atgaaggggg actacgtgct gggggggctg ttccccctgg gcgaggccga ggaggctggc 360
     38 ctccgcagcc ggacacggcc cagcagccct gtgtgcacca ggtacagagg tgggacggcc 420
     39 tgggtcgggg tcagggtgac caggtctggg gtgctcctga gctggggccg aggtggccat 480
    40 ctgcggttct gtgtggcccc aggttctcct caaacggcct gctctgggca ctggccatga 540
     41 aaatggccgt ggaggagatc aacaacaagt cggatctgct gcccgggctg cgcctgggct 600
     42 acgaectett tgatacgtge teggageetg tggtggeeat gaageeeage eteatgttee 660
     43 tggccaagge aggcageege gacategeeg cetaetgeaa etaeaegeag taecageece 720
     44 gtgtgctggc tgtcatcggg ccccactcgt cagagctcgc catggtcacc ggcaagttct 780
     45 teagettett ceteatgeee eagtggggeg cedeceacea teacecacee ceaaceaace 840
                                                                           876
     46 cctgcccgt gggagcccct tgtgtcagga gaatgc
     49 <210> SEQ ID NO: 2
     50 <211> LENGTH: 2687
     51 <212> TYPE: DNA
     52 <213> ORGANISM: Homo sapiens
     54 <400> SEQUENCE: 2
     55 tacatgcacc ccacccagcc ctgccctggg agccctgtgt cagaagatgc tcttggcctt 60
     56 gcaggtcagc tacggtgcta gcatggagct gctgagcgcc cgggagacct tcccctcctt 120
     57 cttccqcacc qtqcccaqcq accqtqtqca gctgacggcc gccgcggagc tgctgcagga 180
```

DATE: 10/18/2004 TIME: 09:54:28

PATENT APPLICATION: US/10/725,284

Input Set : A:\78003681.app

```
58 gttcggctgg aactgggtgg ccgccctggg cagcgacgac gagtacggcc ggcagggcct 240
59 gagcatette teggeeetgg eegeggeaeg eggeatetge ategegeaeg agggeetggt 300
60 geogetgeec egtgeegatg actegegget ggggaaggtg caggaegtee tgeaccaggt 360
61 gaaccagage agegtgeagg tggtgetget gttegeetee gtgeaegeeg ceeaegeeet 420
62 cttcaactac agcatcagca gcaggetete geccaaggtg tgggtggeca gcgaggeetg 480
63 gctgacctct gacctggtca tggggctgcc cggcatggcc cagatgggca cggtgcttgg 540
64 cttcctccaq aqqqqtqccc agctgcacga gttcccccag tacgtgaaga cgcacctggc 600
65 cctggccacc gacccggcct tctgctctgc cctgggcgag agggagcagg gtctggagga 660
66 ggacgtggtg ggccagcgct gcccgcagtg tgactgcatc acgctgcaga acgtgagcgc 720
67 agggctaaat caccaccaga cgttctctgt ctacgcagct gtgtatagcg tggcccaggc 780
68 cctgcacaac actcttcagt gcaacgcctc aggctgcccc gcgcaggacc ccgtgaagcc 840
69 ctggcaggtg agcccgggag atgggggtgt gctgtcctct gcatgtgccc aggccaccag 900
70 gcacggccac cacgcctgag ctggaggtgg ctggcggctc agccccgtcc cccgcccgca 960
71 gctcctggag aacatgtaca acctgacctt ccacgtgggc gggctgccgc tgcggttcga 1020
72 cagcagcgga aacgtggaca tggagtacga cctgaagctg tgggtgtggc agggctcagt 1080
73 gcccaggctc cacgacgtgg gcaggttcaa cggcagcctc aggacagagc gcctgaagat 1140
74 ccgctggcac acgtctgaca accaggtgag gtgagggtgg gtgtgccagg cgtgcccgtg 1200
75 gtagcccccg cggcagggcg cagcctgggg gtgggggccg ttccagtctc ccgtgggcat 1260
76 qcccaqccqa gcaqagccag accccaggcc tgtgcgcaga agcccgtgtc ccggtgctcg 1320
77 cggcagtgcc aggagggcca ggtgcgccgg gtcaaggggt tccactcctg ctgctacgac 1380
78 tgtgtggact gcgaggcggg cagctaccgg caaaacccag gtgagccgcc ttcccggcag 1440
79 gegggggtgg gaacgcagca ggggagggte etgecaagte etgaetetga gaccagagee 1500
80 cacagggtac aagacgaaca cccagcgccc ttctcctctc tcacagacga catcgcctgc 1560
81 accttttgtg gccaggatga gtggtccccg gagcgaagca cacgctgctt ccgccgcagg 1620
82 teteggttee tggeatgggg egageegget gtgetgetge tgeteetget getgageetg 1680
84 gttcaggcct cgggggggcc cctggcctgc tttggcctgg tgtgcctggg cctggtctgc 1800
85 ctcagcgtcc tectgttece tggccagece agecetgece gatgeetgge ccagcagece 1860
86 ttgtcccacc tcccgctcac gggctgcctg agcacactct tcctgcaggc ggccgagatc 1920
87 ttegtqqaqt cagaactgee tetgagetgg geagacegge tgagtggetg cetgeggggg 1980
88 ccctgggcct ggctggtggt gctgctggcc atgctggtgg aggtcgcact gtgcacctgg 2040
89 tacctggtgg cettecegee ggaggtggtg aeggaetgge acatgetgee caeggaggeg 2100
90 ctggtgcact geegeacaeg eteetgggte agetteggee tagegeaege caccaatgee 2160
91 acgctggcct ttctctgctt cctgggcact ttcctggtgc ggagccagcc gggctgctac 2220
92 aaccgtgccc gtggcctcac ctttgccatg ctggcctact tcatcacctg ggtctccttt 2280
93 gtgcccctcc tggccaatgt gcaggtggtc ctcaggcccg ccgtgcagat gggcgccctc 2340
94 ctgctctgtg tcctgggcat cctggctgcc ttccacctgc ccaggtgtta cctgctcatg 2400
95 cggcagccag ggctcaacac ccccgagttc ttcctgggag ggggccctgg ggatgcccaa 2460
96 ggccagaatg acgggaacac aggaaatcag gggaaacatg agtgacccaa ccctgtgatc 2520
97 tcagccccgg tgaacccaga cttagctgcg atccccccca agccagcaat gacccgtgtc 2580
98 tegetacaga gacceteceg etetaggite tgaccecagg tigietectg accetgacce 2640
                                                                    2687
99 cacagtgage cetaggeetg gageacgtgg acaeceetgt gaccate
102 <210> SEQ ID NO: 3
103 <211> LENGTH: 2559
104 <212> TYPE: DNA
105 <213 > ORGANISM: Homo sapiens
107 <400> SEQUENCE: 3
108 atgctgggcc ctgctgtcct gggcctcagc ctctgggctc tcctgcaccc tgggacgggg 60
109 gccccattgt gcctgtcaca gcaacttagg atgaaggggg actacgtgct gggggggctg 120
```

RAW SEQUENCE LISTING DATE: 10/18/2004 PATENT APPLICATION: US/10/725,284 TIME: 09:54:28

Input Set : A:\78003681.app

```
110 ttcccctgg gcgaggccga ggaggctggc ctccgcagcc ggacacggcc cagcagccct 180
111 gtgtgcacca ggttctcctc aaacggcctg ctctgggcac tggccatgaa aatggccgtg 240
112 gaggagatca acaacaagtc ggatctgctg cccgggctgc gcctgggcta cgacctcttt 300
113 gatacgtgct cggagcctgt ggtggccatg aagcccagcc tcatgttcct ggccaaggca 360
114 ggcagccgcg acategeege etactgcaac tacacgcagt accageceeg tgtgctggct 420
115 gtcatcgggc cccactcgtc agagctcgcc atggtcaccg gcaagttctt cagcttcttc 480
116 ctcatgcccc aggtcagcta cggtgctagc atggagctgc tgagcgcccg ggagaccttc 540
117 ccctccttct tccgcaccgt gcccagcgac cgtgtgcagc tgacggccgc cgcggagctg 600
118 ctgcaggagt tcggctggaa ctgggtggcc gccctgggca gcgacgacga gtacggccgg 660
119 cagggeetga geatettete ggeeetggee geggeaegeg geatetgeat egegeaegag 720
120 ggcctggtgc cgctgccccg tgccgatgac tcgcggctgg ggaaggtgca ggacgtcctg 780
121 caccaggtga accagagcag cgtgcaggtg gtgctgctgt tcgcctccgt gcacgccgcc 840
122 cacgccctct tcaactacag catcagcagc aggctctcgc ccaaggtgtg ggtggccagc 900
123 gaggeetgge tgaeetetga eetggteatg gggetgeeeg geatggeeea gatgggeaeg 960
124 qtqcttqqct tcctccaqaq qqqtqcccaq ctqcacgagt tcccccagta cgtgaagacg 1020
125 cacctggccc tggccaccga cccggccttc tgctctgccc tgggcgagag ggagcagggt 1080
126 ctggaggagg acgtggtggg ccagcgctgc ccgcagtgtg actgcatcac gctgcagaac 1140
127 gtgagcgcag ggctaaatca ccaccagacg ttctctgtct acgcagctgt gtatagcgtg 1200
128 gcccaqqccc tqcacaacac tcttcagtgc aacgcctcag gctgccccgc gcaggacccc 1260
129 gtgaagccct ggcagctcct ggagaacatg tacaacctga ccttccacgt gggcgggctg 1320
130 ccgctgcggt tcgacagcag cggaaacgtg gacatggagt acgacctgaa gctgtgggtg 1380
131 tggcagggct cagtgcccag gctccacgac gtgggcaggt tcaacggcag cctcaggaca 1440
132 gagcgcctga agatccgctg gcacacgtct gacaaccaga agcccgtgtc ccggtgctcg 1500
133 eggeagtgee aggagggeea ggtgegeegg gteaaggggt teeaeteetg etgetaegae 1560
134 tgtgtggact gcgaggcggg cagctaccgg caaaacccag acgacatcgc ctgcaccttt 1620
135 tgtggccagg atgagtggtc cccggagcga agcacacgct gcttccgccg caggtctcgg 1680
136 ttcctggcat ggggcgagcc ggctgtgctg ctgctgctcc tgctgctgag cctggcgctg 1740
137 ggccttgtgc tggctgcttt ggggctgttc gttcaccatc gggacagccc actggttcag 1800
138 gcctcggggg ggcccctggc ctgctttggc ctggtgtgcc tgggcctggt ctgcctcagc 1860
139 qtcctcctqt tccctqqcca gcccaqccct gcccgatgcc tggcccagca gcccttgtcc 1920
140 cacctcccgc tcacgggctg cctgagcaca ctcttcctgc aggcggccga gatcttcgtg 1980
141 gagtcagaac tgcctctgag ctgggcagac cggctgagtg gctgcctgcg ggggccctgg 2040
142 gcctggctgg tggtgctgct ggccatgctg gtggaggtcg cactgtgcac ctggtacctg 2100
143 gtggccttcc cgccggaggt ggtgacggac tggcacatgc tgcccacgga ggcgctggtg 2160
144 cactgoogca cacgotoctg ggtcagotto ggcctagogc acgocaccaa tgccacgctg 2220
145 gcctttctct gcttcctggg cactttcctg gtgcggagcc agccgggctg ctacaaccgt 2280
146 gcccgtggcc tcacctttgc catgctggcc tacttcatca cctgggtctc ctttgtgccc 2340
147 ctcctggcca atgtgcaggt ggtcctcagg cccgccgtgc agatgggcgc cctcctgctc 2400
148 tgtgtcctgg gcatcctggc tgccttccac ctgcccaggt gttacctgct catgcggcag 2460
149 ccaqqqctca acacccccqa qttcttcctq ggagqqggcc ctggggatgc ccaaggccag 2520
                                                                      2559
150 aatgacgga acacaggaaa tcaggggaaa catgagtga
153 <210> SEQ ID NO: 4
154 <211> LENGTH: 852
155 <212> TYPE: PRT
156 <213> ORGANISM: Homo sapiens
158 <400> SEQUENCE: 4
159 Met Leu Gly Pro Ala Val Leu Gly Leu Ser Leu Trp Ala Leu Leu His
                                         10
160
162 Pro Gly Thr Gly Ala Pro Leu Cys Leu Ser Gln Gln Leu Arg Met Lys
```

DATE: 10/18/2004

PATENT APPLICATION: US/10/725,284 TIME: 09:54:28

Input Set : A:\78003681.app

163				20					25					30		
	Gly	Asp	Tvr		Leu	Glv	Glv	Leu		Pro	Leu	Glv	Glu		Glu	Glu
166	1		35			1	- 4	40				1	45			
	Ala	Glv		Ara	Ser	Ara	Thr		Pro	Ser	Ser	Pro	Val	Cvs	Thr	Arq
169		50		5		5	55					60		-		~
	Phe		Ser	Asn	Glv	Leu		Trp	Ala	Leu	Ala	Met	Lys	Met	Ala	Val
172	65				- 1	70		_			75		-			80
	Glu	Glu	Ile	Asn	Asn	Lvs	Ser	Asp	Leu	Leu	Pro	Gly	Leu	Arq	Leu	Gly
175					85	-		-		90		•			95	-
	Tyr	Asp	Leu	Phe	Asp	Thr	Cys	Ser	Glu	Pro	Val	Val	Ala	Met	Lys	Pro
178	•	-		100	-		•		105					110	_	
	Ser	Leu	Met	Phe	Leu	Ala	Lys	Ala	Gly	Ser	Arg	Asp	Ile	Ala	Ala	Tyr
181			115				-	120	-		_	_	125			_
183	Cys	Asn	Tyr	Thr	Gln	Tyr	Gln	Pro	Arg	Val	Leu	Ala	Val	Ile	Gly	Pro
184	_	130	-			_	135		_			140				
186	His	Ser	Ser	Glu	Leu	Ala	Met	Val	Thr	Gly	Lys	Phe	Phe	Ser	Phe	Phe
	145					150					155					160
189	Leu	Met	Pro	Gln	Val	Ser	Tyr	Gly	Ala	Ser	Met	Glu	Leu	Leu	Ser	Ala
190					165		_			170					175	
192	Arg	Glu	Thr	Phe	Pro	Ser	Phe	Phe	Arg	Thr	Val	Pro	Ser	Asp	Arg	Val
193	_			180					185					190		
195	Gln	Leu	Thr	Ala	Ala	Ala	Glu	Leu	Leu	Gln	Glu	Phe	Gly	Trp	Asn	$\mathtt{Trp}$
196			195					200					205			
198	Val	Ala	Ala	Leu	Gly	Ser	Asp	Asp	Glu	Tyr	Gly	Arg	Gln	Gly	Leu	Ser
199		210					215					220				
201	Ile	Phe	$\operatorname{\mathtt{Ser}}$	Ala	Leu	Ala	Ala	Ala	Arg	Gly	Ile	Cys	Ile	Ala	His	Glu
	225					230					235					240
204	Gly	Leu	Val	Pro	Leu	Pro	Arg	Ala	Asp	Asp	Ser	Arg	Leu	Gly	Lys	Val
205					245					250					255	
207	Gln	Asp	Val	Leu	His	Gln	Val	Asn		Ser	Ser	Val	Gln		Val	Leu
208				260					265			_		270		
210	Leu	Phe		Ser	Val	His	Ala		His	Ala	Leu	Phe		Tyr	Ser	Ile
211			275				_	280				_	285		_	_
	Ser		Arg	Leu	Ser	Pro		Val	Trp	Val	Ala		GLu	Ala	Trp	Leu
214		290	_	_			295	_	_	~ 7		300	~7		~7	ml
	Thr	Ser	Asp	Leu	vaı		GLY	Leu	Pro	GTA		Ala	GIN	мет	GIY	
	305	_	~-7	-1	_	310	_	~7	~ 7	~7	315	** *	<b>~</b> 1	D1	D	320
	Val	Leu	GLY	Phe		GIn	Arg	GIY	Ala		Leu	HIS	GIU	Pne		GIII
220	_		_	m1.	325	<b>.</b>	<b>33</b> -	<b>T</b>	<b>7</b> . 7	330	3	D	7. T _	Dl	335	C
	Tyr	vai	ьуs		HIS	ьeu	Ara	ьeu		Tnr	Asp	Pro	Ата		Cys	ser
223		_	~7	340		<b>~</b> 1	<b>~1</b>	<b>a</b> 1	345	<b>a</b> 1	<b>~1</b>	7	**- 7	350	<b>~1</b>	C1
	Ala	ьeu		GLU	arg	GIU	GIN		ьeu	GIU	GLU	Asp		val	σιγ	Gln
226	<b>.</b>	<b>a</b> -	355	<b>~</b> 1.	<b>C</b>	<b>7</b>	<b>C</b>	360	m1	т	a1	71	365	0	71 -	C1
	Arg		Pro	GIN	Cys	Asp		тте	Tnr	ьeu	GIN		vaı	ser	Ala	GIY
229	т	370	TT	TT-1	<b>~1</b>	ml	375	Cox	77~ T	m,	η T -	380	¥7 ~ 7	Ф	Cor	77⊃7
	Leu	Asn	HIS	HIS	GIII		rne	ser	VdI	ıyr		AId	val	TAT	per	400
	385	~1	71 T	T 011	п: ~	390	Th~	T 637	C1 ~	C3~	395	7.1.~	C0~	G1 **	Czzc	
	Ala	GTII	ATG	ьeu		ASII	TIIT	neu	GTII		ASII	ATG	SET	<del>с</del> т у	415	110
235					405					410					-17	

DATE: 10/18/2004 TIME: 09:54:28

PATENT APPLICATION: US/10/725,284

Input Set : A:\78003681.app

237 238	Ala	Gln	Asp	Pro 420	Val	Lys	Pro	$\operatorname{Trp}$	Gln 425	Leu	Leu	Glu	Asn	Met 430	Tyr	Asn
240	Leu	Thr	Phe 435		Val	Gly	Gly	Leu 440		Leu	Arg	Phe	Asp 445		Ser	Gly
	Asn			Met	Glu	Tyr			Lys	Leu	Trp			Gln	Gly	Ser
	Val	450 Pro	Arg	Leu	His		455 Val	Gly	Arg	Phe		460 Gly	Ser	Leu	Arg	
	465 Glu	7	T 0	T	<b>T</b> 10	470	Twn	uic	Thr	Cor	475	λan	Gln	Luc	Pro	480 Val
250					485					490					495	
253	Ser			500					505					510		
255 256	Gly	Phe	His 515	Ser	Cys	Суѕ	Tyr	Asp 520	Cys	Val	Asp	Cys	Glu 525		Gly	Ser
	Tyr	Arg 530		Asn	Pro	Asp	Asp 535	Ile	Ala	Cys	Thr	Phe 540	Cys	Gly	Gln	Asp
	Glu		Ser	Pro	Glu	Arg		Thr	Arg	Cys	Phe	-	Arg	Arg	Ser	Arg
262	545	_				550					555					560
	Phe	Leu	Ala	Trp		Glu	Pro	Ala	Val	Leu 570	Leu	Leu	Leu	Leu	Leu 575	Leu
265	Ser	T 011	7 <b>7</b> 7	T 011	565	T 011	37 a 3	T.011	777 =		T.e.u	Glv	T.e.11	Phe		His
268	ser	ьеu	AIA	580	Gry	ьси	vai	шеα	585	пια	пси	Cly	<b>L</b> Cu	590	Val	111.0
	His	Arg	Asp	Ser	Pro	Leu	Val	Gln	Ala	Ser	Gly	Gly	Pro	Leu	Ala	Cys
271			595	•				600					605			
	Phe	_	Leu	Val	Cys	Leu		Leu	Val	Cys	Leu		Val	Leu	Leu	Phe
274	Pro	610	Cl n	Dro	cor	Dro	615	7\ rc	Cvc	T.011	Δla	620	Gln	Pro	Len	Ser
	625	GIA	GIII	PIO	ser	630	АТА	Arg	Cys	пец	635	GIII	0111	110	пси	640
	His	Leu	Pro	Leu	Thr		Cys	Leu	Ser	Thr		Phe	Leu	Gln	Ala	Ala
280					645					650					655	
	Glu	Ile	Phe		Glu	Ser	Glu	Leu		Leu	Ser	Trp	Ala		Arg	Leu
283	Ser	Glar	Carc	660	λκα	Glaz	Dro	Trn	665 Ala	Trn	T.eu	Val	Val	670	Leu	Ala
286	ser	GIY	675	neu	AIG	Gry	FIO	680	AΙα	пр	пси	Val	685	шси	Lou	1114
	Met	Leu		Glu	Val	Ala	Leu	Cys	Thr	Trp	Tyr	Leu	Val	Ala	Phe	Pro
289		690					695					700				_
	Pro	Glu	Val	Val	Thr		Trp	His	Met	Leu		Thr	Glu	Ala	Leu	
	705		_	en1	7	710	m	77. J	0.000	Dho	715	T 011	7 . T . n	цiс	ת ד ת	720 Thr
294 295	His	Cys	Arg	Thr	725	ser	тър	vai	ser	730	СТУ	ьеи	Ата	птъ	735	1111
		Ala	Thr	Leu		Phe	Leu	Cvs	Phe		Glv	Thr	Phe	Leu		Arg
298		****		740				-1 -	745		2			750		_
300	Ser	Gln	Pro	Gly	Cys	Tyr	Asn	Arg	Ala	Arg	Gly	Leu	Thr	Phe	Ala	Met
301			755					760			_		765	_		_
			Tyr	Phe	Ile	Thr		Val	Ser	Phe	Val		Leu	Leu	Ala	Asn
304		770	17 n 7	77~7	Lou	7~~	775 Pro	Δ1 =	T/= T	Gln	Met	780	د ۱ ۵	Len	Len	Leu
	785	GIII	val	val	пеп	790	FLU	міа	val	GIII	795	ОТУ	AIG	Leu	<u> </u>	800
		Val	Leu	Gly	Ile		Ala	Ala	Phe	His		Pro	Arg	Cys	Tyr	Leu
	4 -			-												

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 10/18/2004 TIME: 09:54:29

PATENT APPLICATION: US/10/725,284

Input Set : A:\78003681.app
Output Set: N:\CRF4\09282004\J725284.raw

### Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 3,9,12,18
Seq#:6; N Pos. 3,6,18,21
Seq#:15; N Pos. 1251,1252,1253,1254,1255,1256,1257,1258,1259,1260,1261,1262
Seq#:15; N Pos. 1263,1264,1265,1266,1267,1268,1269,1270,1271,1272,1273,1274
Seq#:15; N Pos. 1275,1276,1277,1278,1279,1280,1281,1282,1283,1284,1285,1286
Seq#:15; N Pos. 1287,1288,1289,1290,1291,1292,1293,1294,1295,1296,1297,1298
Seq#:15; N Pos. 1299,1300,1951,1952,1953,1954,1955,1956,1957,1958,1959,1960
Seq#:15; N Pos. 1961,1962,1963,1964,1965,1966,1967,1968,1969,1970,1971,1972
Seq#:15; N Pos. 1973,1974,1975,1976,1977,1978,1979,1980,1981,1982,1983,1984
Seq#:15; N Pos. 1985,1986,1987,1988,1989,1990,1991,1992,1993,1994,1995,1996
Seq#:15; N Pos. 1997,1998,1999,2000
Seq#:18; Xaa Pos. 1,3,4,6,7,8,11,12,13
Seq#:19; Xaa Pos. 1,3,4,7,9,10,11,13,14,15

#### VERIFICATION SUMMARY

PATENT APPLICATION: US/10/725,284

DATE: 10/18/2004 TIME: 09:54:29

Input Set : A:\78003681.app

Output Set: N:\CRF4\09282004\J725284.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

L:383 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0

L:784 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1200

M:341 Repeated in SeqNo=15

L:1175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0

L:1239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0